

Figure 1:

5	SCS0007 FLJ32009 HCP50656.1	MWAGLLLRACVALLLPAPARGYTGRKPPGHFAAERRRLGPHV MWAGLLLRACVALLLPAPARGYTGRKPPGHFAAERRRLGPHV RGAGAEPSPGDPPFPVMWAGLLLRACVALLLPAPARGYTGRKPPGHFAA RRLGPHV *****
10	SCS0007 FLJ32009 HCP50656.1	CLSGFGSGCCPGWAPSMGGGHCTPLCSFGCGSGICIAPNVCSQDGEQGATCPETHGPC CLSGFGSGCCPGWAPSMGGGHCTPLCSFGCGSGICIAPNVCSQDGEQGAT CPETHGPC CLSGFGSGCCPGWAPSMGGGHCTPLCSFGCGSGICIAPNVCSQDGEQGATCPETHGPC *****
15	SCS0007 FLJ32009 HCP50656.1	GEYGCDLTCNHGGCQEVAVCPVGFSGMTETAVGIRCTDIDECVTSSCEGHCV NTEGGFVC GEYGCDLTCNHGGCQEVAVCPVGFSGMTETAVGIRCTDIDECVTSSCEGHCVNTEGGFVC GEYGCDLTCNHGGCQEVAVCPVGFSGMTETAVGIRCTDIDECVTSSCEGHCVNTEGGFVC *****
20	SCS0007 FLJ32009 HCP50656.1	ECGPGMQLSADRHSCQDTECLGTPCQQRCKNSIGSYKCSRTGFHLHGNRHSCVDVNEC ECGPGMQLSADRHSCQDTECLGTPCQQRCKNSIGSYKCSRTGFHLHGNRHSCVDVNEC ECGPGMQLSADRHSCQDTECLGTPCQQRCKNSIGSYKCSRTGFHLHGNRHSCVDVNEC *****
25	SCS0007 FLJ32009 HCP50656.1	RRPLERRVCHHSCHNTVGSFLCTCRPGFRLRADRVSCGE ----- RRPLERRVCHHSCHNTVGSFLCTCRPGFRLRADRVSCGEAFKAVLAPSAILQP RQHPSKM RRPLERRVCHHSCHNTVGSFLCTCRPGFRLRADRVSCGE ----- *****
30	SCS0007 FLJ32009 HCP50656.1	-----ALSP-----PDWQQGP-----LPAGT----- LLLLPEAGRPALESPGHSFPGSAGFPAGVTRTLPSTPRLPTSSPAPVWLLSTLLATP -----ALSP-----PDWQQGP-----LPAGT----- **** * ** **
35	SCS0007 FLJ32009 HCP50656.1	-----WPCMNQGVAGQSLG----- VPTASLLGNLRPPSLLQGEVMGTPSSPRGPESPRLAAGPSPCWHLGAMHESRSRWTEPGC -----WPCMNQGVAGQSLG----- ** * *
40	SCS0007 FLJ32009 HCP50656.1	-----VPS-AGARLETCTACFVS----- SQCWCEGDGKVTCEKVRCEAACSHPIPSRDGGCCPCTGCFHSGVVRAEGDVFSP PNENCT -----VPS-AGARLETCTACFVS----- ** * * *
45	SCS0007 FLJ32009 HCP50656.1	---VLLAR-----VRPPIKTDCTCVPVRCYFHGRWYADGAVFSGGG DECTTC VCVCLAGNVSCISPECPSPGCPQTPPTDCTCVPVRCYFHGRWYADGAVFSGGGDECTTC ---VLLAR-----VRPPIKTDCTCVPVRCYFHGRWYADGAVFSGGGDECTTC * ** * *****
50	SCS0007 FLJ32009 HCP50656.1	VCQNGEVECSFMPCPELACPREEWRLGPGQCCFTCQEPSTGCSLDDNGVEFFIGQIWS VCQNGEVECSFMPCPELACPREEWRLGPGQCCFTCQEPSTGCSLDDNGVEFFIGQIWS VCQNGEVECSFMPCPELACPREEWRLGPGQCCFTCQEPSTGCSLDDNGVEFFI GOIWS *****
55	SCS0007 FLJ32009 HCP50656.1	PGDPCELCICQADGVSCKRTDCVDSCPHPPIRIPGQCCPDCAAGAQRMLSLAGCTYTGR PGDPCELCICQADGVSCKRTDCVDSCPHPPIRIPGQCCPDCAAG -----CTYTGR PGDPCELCICQADGVSCKRTDCVDSCPHPPIRIPGQCCPDCAAGAQRMLSLAGCTYTGR *****
60	SCS0007 FLJ32009 HCP50656.1	IFYNNETFPSVLDPCLSCICLLGSVACSPVDCPITCTYPFHPDGECCPVCRDCNY EGRKV IFYNNETFPSVLDPCLSCICLLGSVACSPVDCPITCTYPFHPDGECCPVCRDCNYEGRKV IFYNNETFPSVLDPCLSCICLLGSVACSPVDCPITCTYPFHPDGECCPVCRDCNYEGRKV *****

5	SCS0007 FLJ32009 HCP50656.1	ANGQVFTLDDEPCTRCTCQLGEVSCEKVPCQRACADPALLPGDCCSSCPDLSPLEEKQG ANGQVFTLDDEPCTRCTCQLGEVSCEKVPCQRACADPALLPGDCCSSCPDLSPLEEKQG ANGQVFTLDDEPCTRCTCQLGEVSCEKVPCQRACADPAL LPGDCCSSCPDSPQPLQ----- *****
10	SCS0007 FLJ32009 HCP50656.1	LSPHGNVAFSKAGRSLHGDTEAPVNCSSCPGEPPTASPSRPVLHLLQLLLRTNLMKTQTLF LSPHGNVAFSKAGRSLHGDTEAPVNCSSCPGEPPTASPSR PVLHLLQLLLRTNLMKTQTLF --PPPTLAPSSPQWGLLGKSPF ---CCLG-----FPMLHCSFTAYLWEGGKRDPKP * . * * * * * * * *
15	SCS0007 FLJ32009 HCP50656.1	TSPAGAHGPHSLALGLTATFPGEFGASPRLSPGPSTPPG APTLPLASPGAPQPPVTPER TSPAGAHGPHSLALGLTATFPGEFGASPRLSPGPSTPPGAPTLPLASPGAPQPPVTPER TGGKCYFHGRWYTDGAVFSGGGDDCTTCVCQNGEVECSFTPCPELECPRE -----EW * . . * . * . . * . * . *
20	SCS0007 FLJ32009 HCP50656.1	SFSASGAQIVSRWPPLPGTLLTEASALSMDPSPSKTFITLLGPRVLSF -TTSRLSTALA SFSASGAQIVSRWPPLPGTLLTEASALSMDPSPSKTFITLLGPRVLSF -TTSRLSTALA LLGPGQCCFTCREP -----TPTTGCSLDD-----NGVEFPIGQIWSFGDPCRRRLCELQ * . * * * . * . * . * . * *
25	SCS0007 FLJ32009 HCP50656.1	ATTHPGPQQPPVGASRGEESTMLSRFPFAALLIHRLPVGRWET ATTHPGPQQPPVGASRGEESTM ----- EDRLRGLLPAPHPDSWTVLS ----- * * * *

Figure 2:**Sequence****Annotation**

AK056571

weakly similar to FIBULIN -1, ISOFORM A Precursor

XP_166226.

similar to cysteine-rich motor neuron 1; cysteine-rich repeat-containing protein

5

S52 precursor; von Willebrand factor type C domain.

10

SCS0007 MWAGLLLRRAA CVALLLPGAP ARGYTGRKPP GHFAAERRRL GPHVCLSGFG
 AK056571 MWAGLLLRRAA CVALLLPGAP ARGYTGRKPP GHFAAERRRL GPHVCLSGFG
 XP_166226. MLLLLP... ..EAGR...

15

SCS0007 SGCCPGWAPS MGGGHCTLPL CSFGCGSGIC IAPNVCSCQD GEQGATCPET
 AK056571 SGCCPGWAPS MGGGHCTLPL CSFGCGSGIC IAPNVCSCQD GEQGATCPET
 XP_166226. PA LSPGHS.... .. PPS

20

SCS0007 HGPCGEYGCD LTCNHGGCQE VARVCPVGFS MTETAVGIRC TDIDECVTSS
 AK056571 HGPCGEYGCD LTCNHGGCQE VARVCPVGFS MTEAAVGIRC TDIDECVTSS
 XP_166226. GAP..... ..GPP... ..AGVRT T.....

25

SCS0007 CEGHCVNTEG GFVCECGPGM QLSADRHSCQ DTDECLGTPC QQRCKNSIGS
 AK056571 CEGHCVNTEG GFVCECGPGM QLSADRHSCQ DTDECLGTPC QQRCKNSIGS
 XP_166226.RLPSPPT PRLTSSP..

30

SCS0007 YKCSCRTGFH LHGNRHSCVD VNECR RPLER RVCHHSCHNT VGSFLCTCRP
 AK056571 YKCSCRTGFH LHGNRHSCVD VNECR RPLER RVCHHSCHNT VGSFLCTCRP
 XP_166226.SAPVWL LSTLLATPVP TASLLGNLRP

35

SCS0007 GFRLRADRVs CEG..... ..ALSP..
 AK056571 GFRLRADRVs CEAFPKAVLA PSAILQPRQH PSKMLLLLPE AGRPALSPPGH
 XP_166226. PSLIQGEVMG TPS..... ..SP..

40

SCS0007 ..PDWQGGP.LPAGT.
 AK056571 SPPSGAPGPP AGVRTTRLPS PTPRLPTSS P SAPVWLLSTL LATPVPTASL
 XP_166226. ...RGPESP.RLAAGP.

45

SCS0007WPCMNQG VAGQSLG...
 AK056571 LGNLRPPSLL QGEVMGTPSS PRGPESPRLA AGPSPCWHLG AMHESRSRWT
 XP_166226.SPCWHLG AMHESRSRWT

50

SCS0007 VP S.AGARLETC RACFVS....
 AK056571 EPGCSQCWCE DGKVTCEKVR CEAACSHPIP SRDGGCCPSC TGCFFHSGVVR
 XP_166226. EPGCSQCWCE DGKVTCEKVR CEAACSHPIP SR DGGCCPSC TGCFFHSGVVR

55

SCS0007VLLA R.....VRPPIK TDCCTCVFVR
 AK056571 AEGDVFSPFN ENCTVCVCLA GNVSCISPEC PSGPCQTPPQ TDCCTCVFVR
 XP_166226. AEGDVFSPFN ENCTVCVCLA GNVSCISPEC PSGPCQTPPQ TDCCTCVFVR

5 SCS0007 CYFHGRWYAD GAVFSGGGDE CTTCVCQNGE VECSEFMPCPE LACPREEWRL
AK056571 CYFHGRWYAD GAVFSGGGDE CTTCVCQNGE VECSEFMPCPE LACPREEWRL
XP_166226. CYFHGRWYAD GAVFSGGGDE CTTCVCQNGE VECSEFMPCPE LACPREEWRL

10 SCS0007 GPGQCCFTCQ EPTPSTGCSL DDNGVEFFIG QIWS PGDPCE LCICQADGSV
AK056571 GPGQCCFTCQ EPTPSTGCSL DDNGVEFFIG QIWS PGDPCE LCICQADGSV
XP_166226. GPGQCCFTCQ EPTPSTGCSL DDNGVEFFIG QIWS PGDPCE LCICQADGSV

15 SCS0007 SCKRTDCVDS CPHPIRIPGQ CCPDCSAAGA QRMLSLAGCT YTGRIFYNNE
AK056571 SCK RTDCVDS CPHPIRIPGQ CCPDCSAG..CT YTGRIFYNNE
XP_166226. SCKRTDCVDS CPHPIRIPGQ CCPDCSAG..CT YTGRIFYNNE

20 SCS0007 TFPSVLDPCL SCICLLGSA CSPVDCPITC TYFFHPDGEC CPVCRDCNYE
AK056571 TFPSVLDPCL SCICLLGSA CSPVDCPITC TYFFHPDG EC CPVCRDCNYE
XP_166226. TFPSVLDPCL SCICLLGSA CSPVDCPITC TYFFHPDGEC CPVCRDCNYE

25 SCS0007 GRKVANGQVF TLDDEPCTRC TCQLGEVSCE KVPCQRACAD PALLPGDCCS
AK056571 GRKVANGQVF TLDDEPCTRC TCQLGEVSCE KVPCQRACAD PALLPGDCCS
XP_166226. GRKVANG QVF TLDDEPCTRC TCQLGEVSCE KVPCQRACAD PALLPGDCCS

30 SCS0007 SCPDLSPL EKGGLSPHGN VAFSKAGRSL HGDTEAPVNC SSCPGPPTAS
AK056571 SCPDLSPL EKGGLSPHGN VAFSKAGRSL HGDTEAPVNC SSCPGPPTAS
XP_166226. SCPDLSPL EKGGLSPHGN VAFSKAGRSL HGDTEAPVNC S SCPGPPTAS

35 SCS0007 PSRPVLHLLQ LLLRTNLMKT QTLPTSPAGA HGPHSLALGL TATFPGEFGA
AK056571 PSRPVLHLLQ LLLRTNLMKT QTLPTSPAGA HGPHSLALGL TATFPGEFGA
XP_166226. PSRPVLHLLQ LLLRTNLMKT QTLPTSPAGA HGPHSLALGL TATFPGEFGA

40 SCS0007 SPRLSPGPS T PPGAPTLP LA SPGAPQPPV TPERSFSASG AQIVSRWPPL
AK056571 SPRLSPGPST PPGAPTLP LA SPGAPQPPV TPERSFSASG AQIVSRWPPL
XP_166226. SPRLSPGPST PPGAPTLP LA SPGAPQPPV TPERSFSASG AQIVSRWPPL

45 SCS0007 PGTLLEASA LSMDPSPSK TPITLLGPRV LSPTTSRLST ALA ATTHPGP
AK056571 PGTLLEASA LSMDPSPSK TPITLLGPRV LSPTTSRLST ALAATTHPGP
XP_166226. PGTLLEASA LSMDPSPSK TPITLLGPRV LSPTTSRLST ALAATTHPGP

50 SCS0007 QQPPVGASRG EESTMLSRFP HAALLIHRP VGRWET
AK056571 QQPPVGASRG EESTM.....
XP_166226. QQPPVGASRG EESTM.....